

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Sidransky, David
Baylin, Stephen
- (ii) TITLE OF INVENTION: METHOD FOR DETECTION OF NEOPLASTIC CELLS
- (iii) NUMBER OF SEQUENCES: 13
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Fish & Richardson P.C.
 - (B) STREET: 4225 Executive Square, Suite 1400
 - (C) CITY: La Jolla
 - (D) STATE: CA
 - (E) COUNTRY: USA
 - (F) ZIP: 92037
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US
 - (B) FILING DATE: 30-JUNE-1995
 - (C) CLASSIFICATION:
- (vii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Haile Ph.D, Lisa A.
 - (B) REGISTRATION NUMBER: 38,347
 - (C) REFERENCE/DOCKET NUMBER: 07265/061001
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 619/678-5070
 - (B) TELEFAX: 619/678-5099

(2) INFORMATION FOR SEQ ID NO:1:

- (1) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 780 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (11) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:1:

TCCCGAGGCA GTTATGTGAA ATATGGCCTC GATCTTGGAG GTCCGGGTGG GAGTGGGGGT	60
GGGGTGGGGG TGGGGGTGAA GGTGGGGGGC GGGCGCGCTC AGGGAAGGCG GGTGCGCGCC	120
TGCGGGGCGG AGATGGGCAG GGGGCGGTGC GTGGGTCCCA GTCTGCAGTT AAGGGGGCAG	180
GAGTGGCGCT GGTCACTCTT GGTGCCAAG GCGGCGCAG CGGCTGCCGA GTCGGCCCT	240
GGAGGCGGCG AGAACATGGT GCGCAGGTC TTGGTGACCC TCCGGATTCC GCGCGCGTGC	300
GGCCCCCGCG GAGTGAGGGT TTTCGTGGTT CACATCCCGC GGCTCACGGG GGAGTGGGCA	360
GCGCCAGGGG CGCCCGCCGC TGTGGCCCTC GTGCTGATGC TACTGAGGAG CCAGCGTCTA	420
GGGCAGCAGC CGCTTCCTAG AAGACCAGGT AGGAAAGGCC CTCGAAAAGT CCGGGGCGCA	480
CTTGTITTTGT TTGGTGTGTG ATTTCTGAAA CAGATAATTC GTCTCTAGCC CATTCTAGGA	540
GGAGGAGGAG ATAACCGCGG TGGAGGCTTC CCATTGGGT TACAACGACT TAGACATGTG	600
GTCTCGCAG TACCATTGAA CCTGGACCTC CCTTCACACA GCCCTCAATC GTGGGAAACT	660
GAGGCGAACA GAGCTTCTAA ACCCACCTCA GAATCAGTG AGTCCCGAAT ATCCTGGGTG	720
GGAATGACTA AGACACACAC ACACACACAC ACACACACAC ACACACACAG TAGGAAATGT	780

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TCCCGAGGTT TCTCAGAG	18
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(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 30 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: RNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:3:

CAUCAUCAUC AUGATGTCGC ACGGTACCTG

30

(2) INFORMATION FOR SEQ ID NO:4:

(1) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: RNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CUACUACUAC UAACGGGTCG GGTGAGAGTG

30

(2) INFORMATION FOR SEQ ID NO:5:

(1) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:5:

AGTGGCGCTG CTCACCTC

18

(2) INFORMATION FOR SEQ ID NO:6:

(1) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:6:

TCCCGAGGTT TCTCAGAG

18

(2) INFORMATION FOR SEQ ID NO:7:

- (1) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(11) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:7:

GGGTGGGAAA TTGGGTAAG

19

(2) INFORMATION FOR SEQ ID NO:8:

- (1) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(11) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:8:

GGGTCACCAA GAACCTGC

18

(2) INFORMATION FOR SEQ ID NO:9:

- (1) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(11) MOLECULE TYPE: DNA (genomic)

(x1) SEQUENCE DESCRIPTION: SEQ ID NO:9:

TCCCAGTCTG CAGTTAAGG

19

(2) INFORMATION FOR SEQ ID NO:10:

- (1) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(11) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

GTCTAAGTCG TTGTAACCCG

20

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

AGTGCATCAG CACGAGGG

19

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

AACAIGGTGC GCAGGTTG

18

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 36 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GGATCCTAAT ACGACTCACT ATAGGGAGAC CACCAIGGCG CTGCTCACCT CTGGTG

56